OIPE

RAW SEQUENCE LISTING   DATE: 08/28/2001
PATENT APPLICATION: US/09/924,340   TIME: 13:01:08

Input Set : **D:\Seqlist.txt**
Output Set: **N:\CRF3\08282001\I924340.raw**

Does Not Comply
Corrected Diskette Needed

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   2 <110> APPLICANT: Bejanin, Stephane
   3         Tanaka, Hiroaki
   5 <120> TITLE OF INVENTION: HUMAN CDNAS AND PROTEINS AND USES THEREOF
   7 <130> FILE REFERENCE: 91.US2.REG
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/924,340
  10 <141> CURRENT FILING DATE: 2001-08-06
  12 <150> PRIOR APPLICATION NUMBER: US 60/305,456
  13 <151> PRIOR FILING DATE: 2001-07-13
  15 <150> PRIOR APPLICATION NUMBER: US 60/302,277
  16 <151> PRIOR FILING DATE: 2001-06-29
  18 <150> PRIOR APPLICATION NUMBER: US 60/298,698
  19 <151> PRIOR FILING DATE: 2001-06-15
  21 <150> PRIOR APPLICATION NUMBER: US 60/293,574
  22 <151> PRIOR FILING DATE: 2001-05-25
  24 <160> NUMBER OF SEQ ID NOS: 112
  26 <170> SOFTWARE: JPatent
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ERRORED SEQUENCES

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6595 <210> SEQ ID NO: 112
6596 <211> LENGTH: 387
6597 <212> TYPE: PRT
6598 <213> ORGANISM: Homo sapiens
6600 <400> SEQUENCE: 112
6601 Met Ser Thr Phe Phe Ser Asp Thr Ala Trp Ile Cys Leu Ala Val Pro
6602 1               5                  10                 15
6603 Thr Val Leu Cys Gly Thr Val Phe Cys Lys Tyr Lys Lys Ser Ser Gly
6604             20                  25                 30
6605 Gln Leu Trp Ser Trp Met Val Cys Leu Ala Gly Leu Cys Ala Val Cys
6606         35                  40                 45
6607 Leu Leu Ile Leu Ser Pro Phe Trp Gly Leu Ile Leu Phe Ser Val Ser
6608     50                  55                 60
6609 Cys Phe Leu Met Tyr Thr Tyr Leu Ser Gly Gln Glu Leu Leu Pro Val
6610 65                  70                  75                 80
6611 Asp Gln Lys Ala Val Leu Val Thr Gly Gly Asp Cys Gly Leu Gly His
6612             85                  90                 95
6613 Ala Leu Cys Lys Tyr Leu Asp Glu Leu Gly Phe Thr Val Phe Ala Gly
6614             100                 105                110
6615 Val Leu Asn Glu Asn Gly Pro Gly Ala Glu Glu Leu Arg Arg Thr Cys
6616         115                 120                 125
6617 Ser Pro Arg Leu Ser Val Leu Gln Met Asp Ile Thr Lys Pro Val Gln
6618     130                 135                 140
6619 Ile Lys Asp Ala Tyr Ser Lys Val Ala Ala Met Leu Gln Asp Arg Gly
6620 145                 150                 155                160
6621 Leu Trp Ala Val Ile Asn Asn Ala Gly Val Leu Gly Phe Pro Thr Asp
6622             165                 170                175
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6623  Gly Glu Leu Leu Leu Met Thr Asp Tyr Lys Gln Cys Met Ala Val Asn
6624          180                 185                 190
6625  Phe Phe Gly Thr Val Glu Val Thr Lys Thr Phe Leu Pro Leu Leu Arg
6626          195                 200                 205
6627  Lys Ser Lys Gly Arg Leu Val Asn Val Ser Ser Met Gly Gly Gly Ala
6628          210                 215                 220
6629  Pro Val Glu Arg Leu Ala Ser Tyr Gly Ser Ser Lys Ala Ala Val Thr
6630  225                 230                 235                 240
6631  Met Phe Ser Ser Val Met Arg Leu Glu Leu Ser Lys Trp Gly Ile Lys
6632                  245                 250                 255
6633  Val Ala Ser Ile Gln Pro Gly Gly Phe Leu Thr Asn Ile Ala Gly Thr
6634          260                 265                 270
6635  Ser Asp Lys Trp Glu Lys Leu Glu Lys Asp Ile Leu Asp His Leu Pro
6636          275                 280                 285
6637  Ala Glu Val Gln Glu Asp Tyr Cys Gln Asp Tyr Ile Leu Ala Gln Arg
6638          290                 295                 300
6639  Asn Phe Leu Leu Leu Ile Asn Ser Leu Ala Ser Lys Asp Phe Ser Pro
6640  305                 310                 315                 320
6641  Val Leu Arg Asp Ile Gln His Ala Ile Leu Ala Lys Ser Pro Phe Ala
6642                  325                 330                 335
6643  Tyr Tyr Thr Pro Gly Lys Gly Ala Tyr Leu Trp Ile Cys Leu Ala His
6644          340                 345                 350
6645  Tyr Leu Pro Ile Gly Ile Tyr Asp Tyr Phe Ala Lys Arg His Phe Gly
6646          355                 360                 365
6647  Gln Asp Lys Pro Met Pro Arg Ala Leu Arg Met Pro Asn Tyr Lys Lys
6648          370                 375                 380
6649  Lys Ala Pro
6650  385
E--> 6651 (106)
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VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/924,340**

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TIME: 13:01:10

Input Set : **D:\Seqlist.txt**
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L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:1470 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27
L:1470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1491 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27
L:1491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1537 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1551 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1592 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:29
L:1592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1655 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:4089 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:71
L:4089 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:4156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72
L:6651 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:112
```